# UBE2Z (USE1) [6His-tagged] E2 - FAT10 or Ubiquitin Conjugating Enzyme

Alternate Names: UBA6-Specific E2: USE1

Cat. No.	62-0086-100
Lot. No.	1837

Quantity: 100 µg -70°C

FOR RESEARCH USE ONLY

The enzymes of the FATylation path-

way play a pivotal role in a number of

cellular processes including regulated

and targeted proteasomal degradation

of ubiquitylated substrate proteins (Bu-

chsbaum et al., 2011). Three classes

of enzymes are involved in the process

of FATylation; the activating enzyme

Uba6, conjugating enzymes (E2s) and

protein ligases (E3s). UBE2Z is a mem-

ber of the E2 conjugating enzyme fam-

ily and cloning of the human gene was first described by Gu et al. (2007) and

Jin et al. (2007). UBE2Z is widely ex-

pressed in human tissues and expres-

sion is particularly high in the placenta,

pancreas, spleen and testis (Gu et al., 2007). UBE2Z has been identified as an interaction partner of FAT10. FAT10

can be transferred from Uba6 to UBE2Z

in vitro and both FAT10 and UBE2Z

have been co-immunoprecipitated from

intact cells. Down regulation of UBE2Z

by siRNA resulted in a strong reduction

of endogenous conjugate formation

suggesting UBE2Z is the major E2 con-

jugating enzyme in the FAT10 cascade

Aichem, A., C. Pelzer, et al. (2010) USE1 is a bispecific conjugating enzyme for ubiquitin and FAT10, which FAT10ylates itself

Buchsbaum, S., B. Bercovich, et al. (2011) FAT10 is a proteasomal degradation signal which is itself regulated by ubiquitina-

Gu, X., F. Zhao, et al. (2007) Cloning and characterization of

a gene encoding the human putative ubiquitin conjugating en-

Jin, J., X. Li, et al. (2007) Dual E1 activation systems for

ubiquitin differentially regulate E2 enzyme charging. Nature

zyme E2Z (UBE2Z). Mol Biol Rep 34(3): 183-8.

(Aichem et al., 2010).

**References:** 

447(7148): 1135-8.

in cis. Nat Commun 1: 13.

tion. Mol Biol Cell. (in press)

Background

**Physical Characteristics** 

Species: human

Source: E. coli expression

Quantity: 100 µg

Concentration: 1 mg/ml

Formulation: 50 mM HEPES pH 7.5. 150 mM sodium chloride, 2 mM dithiothreitol, 10% glycerol

Molecular Weight: ~40.6 kDa

Purity: >90% by InstantBlue™ SDS-PAGE

Stability/Storage: 12 months at -70°C; aliquot as required

# **Quality Assurance**

#### **Purity:**

4-12% gradient SDS-PAGE InstantBlue<sup>™</sup> staining Lane 1: MW markers Lane 2: 1 µg His-UBE2Z





# **Protein Sequence:**

MGSSHHHHHHSSGLEVLFQGPGSMAESPTEE AATAGAGAAGPGASSVAGVVGVSGSGGGFGP PFLPDVWAAAAAAGGAGGPGSGLAPLPGLPP SAAAHGAALLSHWDPTLSSDWDGERTAPOCLL RIKRDIMSIYKEPPPGMFVVPDTVDMTKIHALIT GPFDTPYEGGFFLFVFRCPPDYPIHPPRVKLMTT **GNNTVRFNPNFYRNGKVCLSILGTWTGPAWSPAQ** SISSVLISIQSLMTENPYHNEPGFEQERHPGD SKNYNECIRHETIRVAVCDMMEGKCPCPEPL RGVMEKSFLEYYDFYEVACKDRLHLQGQTMQDPF GEKRGHFDYQSLLMRLGLIRQKVLERLHNENAE MDSDSSSSGTETDLHGSLRV

#### Tag (bold text): N-terminal His

Protease cleavage site: PreScission™ (LEVLFQ▼GP) UBE2Z (regular text): Start bold italics (amino acid residues 1-354) Accession number: NP 075567.2

## **Protein Identification:**

Confirmed by mass spectrometry.

### E2-Ubiquitin Thioester Loading Assay:

The activity of His-UBE2Z was validated by loading E1 Uba6 activated ubiquitin onto the active cysteine of the His-UBE2Z E2 enzyme via a transthiolation reaction. Incubation of the Uba6 and His-UBE2Z enzymes in the presence of ubiquitin and ATP at 30°C was compared at two time points,  $T_0$  and  $T_{10}$  minutes. Under these conditions tested no His-UBE2Z/ ubiquitin thioester loading was observed.

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Lot-specific COA version tracker: v1.0.0

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Storage: NOT FOR USE IN HUMANS